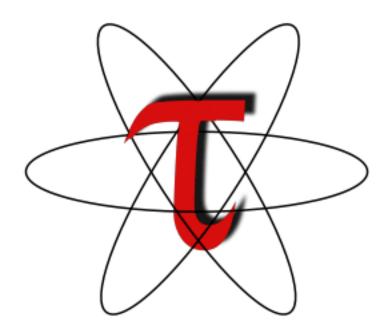
TAU User Guide



TAU User Guide

Updated November 10th 2011 for use with version 2.21 or greater.

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TAU Performance System® is a portable profiling and tracing toolkit for performance analysis of parallel programs written in Fortran, C, C++, Java, and Python. TAU (Tuning and Analysis Utilities) is capable of gathering performance information through instrumentation of functions, methods, basic blocks, and statements. The TAU API also provides selection of profiling groups for organizing and controlling instrumentation. Calls to the TAU API are made by probes inserted into the execution of the application via source transformation, compiler directives or by library interposition.

This guide is organized into different sections. Readers wanting to get started right way can skip to the Common Profile Requests section for step-by-step instructions for obtaining difference kinds of performance data. Or browse the starters guide for a quick reference to common TAU commands and variables.

TAU can be found on the web at: http://tau.uoregon.edu

Chapter 1. Tau Instrumentation

TAU provides three methods to track the performance of your application. Library interposition using tau_exec, compiler directives or source transformation using PDT. Here is a table that lists the features/requirement for each method:

Table 1.1. Different methods of instrumenting applications

Method	Requires recompil- ing	Requires PDT	Shows MPI events	Routine- level event	Low level events (loops, phases, etc)		Ability to exclude file from instrumentation
Interposi- tion			Yes			Yes	
Compiler	Yes		Yes	Yes		Yes	Yes
Source	Yes	Yes	Yes	Yes	Yes	Yes	Yes

The requirements for each method increases as we move down the table: tau_exec only requires a system with shared library support. Compiler based instrumentation requires re-compiling that target application and Source instrumentation aditionally requires PDT. For this reason we often recommend that users start with Library interposition and move down the table if more features are needed.

1.1. Dynamic instrumentation through library pre-loading

Dynamic instrumentation is achieved through library pre-loading. The libraries chosen for pre-loading determine the scope of instrumentation. Some options include tracking MPI, io, memory, cuda, opencl library calls. MPI instrumentation is included by default the others are enabled by command-line options to tau_exec. More info at the tau_exec manual page. Dynamic instrumentation can be used on both uninstrumented binaries and binaries instrumented via one of the methods below, in this way different layers of instrumentation can be combined.

To use tau_exec place this command before the application executable when running the application. In this example IO instrumentation is requested.

```
%> tau_exec -io ./a.out
%> mpirun -np 4 tau_exec -io ./a.out
```

1.2. TAU scripted compilation

For more detailed profiles, TAU provides two means to compile your application with TAU: through your compiler or through source transformation using PDT.

1.2.1. Compiler Based Instrumentation

TAU provides these scripts: tau_f90.sh, tau_cc.sh, and tau_cxx.sh to instrument and compile Fortran, C, and C++ programs respectively. You might use tau cc.sh to compile a C program by typing:

```
%> module load tau
%> tau_cc.sh -tau_options=-optCompInst samplecprogram.c
```

On machines where a TAU module is not available, you will need to set the tau makefile and/or options. The makefile and options controls how will TAU will compile you application. Use

The Makefile can be found in the /[arch]/lib directory of your TAU distribution, for example / x86_64/lib/Makefile.tau-mpi-pdt.

You can also use a Makefile specified in an environment variable. To run tau_cc.sh so it uses the Makefile specified by environment variable TAU_MAKEFILE, type:

```
%>export TAU_MAKEFILE=[path to tau]/[arch]/lib/[makefile]
%>export TAU_OPTIONS=-optCompInst
%>tau_cc.sh sampleCprogram.c
```

Similarly, if you want to set compile time options like selective instrumentation you can use the TAU_OPTIONS environment variable.

1.2.2. Source Based Instrumentation

TAU provides these scripts: tau_f90.sh, tau_cc.sh, and tau_cxx.sh to instrument and compile Fortran, C, and C++ programs respectively. You might use tau_cc.sh to compile a C program by typing:

```
%> module load tau
%> tau_cc.sh samplecprogram.c
```

When setting the TAU_MAKEFILE make sure the Makefile name contains pdt because you will need a version of TAU built with PDT.

A list of options for the TAU compiler scripts can be found by typing man tau_compiler.sh or in this chapter of the reference guide.

1.2.3. Options to TAU compiler scripts

These are some commonly used options available to the TAU compiler scripts. Either set them via the TAU_OPTIONS environment variable or the -tau_options= option to tau_f90.sh, tau_cc.sh, or tau_cxx.sh

```
-optVerbose Enable verbose output (default: on)
-optKeepFiles Do not remove intermediate files
-optShared Use shared library of TAU (consider when using tau_exec
```

1.3. Selectively Profiling an Application

1.3.1. Custom Profiling

TAU allows you to customize the instrumentation of a program by using a selective instrumentation file. This instrumentation file is used to manually control which parts of the application are profiled and how they are profiled. If you are using one of the TAU compiler wrapper scripts to instrument your application you can use the -tau_options=-optTauSelectFile=<file> option to enable selective instrumentation.



Note

Selective instrumentation is only available when using source-level instrumentation (PDT).

To specify a selective instrumentation file, create a text file and use the following guide to fill it in:

- Wildcards for routine names are specified with the # mark (because * symbols show up in routine signatures.) The # mark is unfortunately the comment character as well, so to specify a leading wildcard, place the entry in quotes.
- Wildcards for file names are specified with * symbols.

```
Here is a example file:
#Tell tau to not profile these functions
BEGIN_EXCLUDE_LIST
void quicksort(int *, int, int)
# The next line excludes all functions beginning with "sort_" and having
# arguments "int *'
void sort #(int *)
void interchange(int *, int *)
END_EXCLUDE_LIST
#Exclude these files from profiling
BEGIN FILE EXCLUDE LIST
*.so
END_FILE_EXCLUDE_LIST
BEGIN_INSTRUMENT_SECTION
# A dynamic phase will break up the profile into phase where
# each events is recorded according to what phase of the application
# in which it occured.
dynamic phase name="fool_bar" file="foo.c" line=26 to line=27
# instrument all the outer loops in this routine
loops file="loop_test.cpp" routine="multiply"
# tracks memory allocations/deallocations as well as potential leaks
memory file="foo.f90" routine="INIT"
# tracks the size of read, write and print statements in this routine
io file="foo.f90" routine="RINB"
```

END_INSTRUMENT_SECTION

Selective instrumentation files can be created automatically from ParaProf by right clicking on a trial and selecting the Create Selective Instrumentation File menu item.

Chapter 2. Profiling

This chapter describes running an instrumented application, generating profile data and analyzing that data. Profiling shows the summary statistics of performance metrics that characterize application performance behavior. Examples of performance metrics are the CPU time associated with a routine, the count of the secondary data cache misses associated with a group of statements, the number of times a routine executes, etc.

2.1. Running the Application

After instrumentation and compilation are completed, the profiled application is run to generate the profile data files. These files can be stored in a directory specified by the environment variable PROFILEDIR. By default, profiles are placed in the current directory. You can also set the TAU_VERBOSE environment variable to see the steps the TAU measurement systems takes when your application is running. Example:

```
% setenv TAU_VERBOSE 1
% setenv PROFILEDIR /home/sameer/profiledata/experiment55
% mpirun -np 4 matrix
```

Other environment variables you can set to enable these advanced MPI measurement features are TAU_TRACK_MESSAGE to track MPI message statistics when profiling or messages lines when tracing, and TAU_COMM_MATRIX to generate MPI communication matrix data.

2.2. Reducing Performance Overhead with TAU_THROTTLE

TAU automatically throttles short running functions in an effort to reduce the amount of overhead associated with profiles of such functions. This feature may be turned off by setting the environment variable TAU_THROTTLE to 0. The default rules TAU uses to determine which functions to throttle is: numc-alls > 100000 && usecs/call < 10 which means that if a function executes more than 100000 times and has an inclusive time per call of less than 10 microseconds, then profiling of that function will be disabled after that threshold is reached. To change the values of numcalls and usecs/call the user may optionally set environment variables:

```
% setenv TAU_THROTTLE_NUMCALLS 2000000
% setenv TAU_THROTTLE_PERCALL 5
```

The changes the values to 2 million and 5 microseconds per call. Functions that are throttled are marked explicitly in there names as THROTTLED.

2.3. Profiling each event callpath

You can enable callpath profiling by setting the environment variable TAU_CALLPATH. In this mode TAU will recorded the each event callpath to the depth set by the TAU_CALLPATH_DEPTH environment variable (default is two). Because instrumentation overhead will increase with the depth of the callpath, you should use the shortest call path that is sufficient.

2.4. Using Hardware Counters for Measurement

Performance counters exist on many modern microprocessors. They can count hardware performance events such as cache misses, floating point operations, etc. while the program executes on the processor. The Performance Data Standard and API (PAPI [http://icl.cs.utk.edu/papi/]) package provides a uniform interface to access these performance counters.

To use these counters, you must first find out which PAPI events your system supports. To do so type:

```
%> papi_avail
Available events and hardware information.
Vendor string and code : AuthenticAMD (2)
Model string and code : AMD K8 Revision C (15)
CPU Revision : 2.000000
                          : 2592.695068
CPU Megahertz
CPU's in this Node
Nodes in this System
                        : 1
Total CPU's
Number Hardware Counters: 4
Max Multiplex Counters : 32
The following correspond to fields in the PAPI_event_info_t structure.
Name
                Code
                                          Deriv
                                                  Description (Note)
PAPI_L1_DCM
                0x80000000
                                 Yes
                                          Yes
Yes
                                                  Level 1 data cache misses
PAPI_L1_ICM
               0x80000001
                                                 Level 1 instruction cache misses
                                 Yes
. . .
Next, to test the compatibility between each metric you wish papi to profile, use
papi_event_chooser:
papi/utils> papi_event_chooser PAPI_LD_INS PAPI_SR_INS PAPI_L1_DCM
Test case eventChooser: Available events which can be added with given
events.
         _____
Vendor string and code : GenuineIntel (1)
Model string and code : Itanium 2 (2)
CPU Revision : 1.000000
CPU Megahertz : 1500.000000
CPU's in this Node : 16
Nodes in this System : 1
Nodes in this System
Total CPU's
Number Hardware Counters: 4
Max Multiplex Counters : 32
Event PAPI L1 DCM can't be counted with others
```

Here the event chooser tells us that PAPI_LD_INS, PAPI_SR_INS, and PAPI_L1_DCM are incompatible metrics. Let try again this time removing PAPI_L1_DCM:

```
Model string and code
                         : Itanium 2 (2)
                         : 1.000000
CPU Revision
CPU Megahertz
                         : 1500.000000
CPU's in this Node
                         : 16
                         : 1
Nodes in this System
                         : 16
Total CPU's
Number Hardware Counters: 4
Max Multiplex Counters
                         : 32
Usage: eventChooser NATIVE|PRESET evt1 evet2 ...
```

Here the event chooser verifies that PAPI_LD_INS and PAPI_SR_INS are compatible metrics.

Next, make sure that you are using a makefile with papi in its name. Then set the environment variable TAU_METRICS to a colon delimited list of PAPI metrics you would like to use.

```
setenv TAU_METRICS PAPI_FP_OPS\:PAPI_L1_DCM
```

In addition to PAPI counters, we support TIME (via unix gettimeofday). On Linux and CrayCNL systems, we provide the high resolution LINUXTIMERS metric and on BGL/BGP systems we provide BGLTIMERS and BGPTIMERS.

Chapter 3. Tracing

Typically, profiling shows the distribution of execution time across routines. It can show the code locations associated with specific bottlenecks, but it can not show the temporal aspect of performance variations. Tracing the execution of a parallel program shows when and where an event occurred, in terms of the process that executed it and the location in the source code. This chapter discusses how TAU can be used to generate event traces.

3.1. Generating Event Traces

To enable tracing with TAU, set the environment variable TAU_TRACE to 1. Similarly you can enable/disable profile with the TAU_PROFILE variable. Just like with profiling, you can set the output directory with a environment variable:

% setenv TRACEDIR /users/sameer/tracedata/experiment56

This will generate a trace file and an event file for each processor. To merge these files, use the tau_treemerge.pl script. If you want to convert TAU trace file into another format use the tau2otf, tau2vtf, or tau2slog2 scripts.

Chapter 4. Analyzing Parallel Applications

4.1. Text summary

For a quick view summary of TAU performance, use pprof It reads and prints a summary of the TAU data in the current directory. For performance data with multiple metrics, move into one of the directories to get information about that metric:

```
%> cd MULTI__P_WALL_CLOCK_TIME
%> pprof
Reading Profile files in profile.*
```

NODE 0; CONTEXT 0; THREAD 0:

%Time	Exclusive msec	Inclusive total msec	#Call	#Subrs	Inclusive usec/call	Name
100.0	24	590	1	1	590963	main
95.9	26	566	1	2	566911	multiply
47.3	279	279	1	0	279280	multiply-opt
44.1	260	260	1	0	260860	multiply-regula

4.2. ParaProf

To launch ParaProf, execute paraprof from the command line where the profiles are located. Launching ParaProf will bring up the manager window and a window displaying the profile data as shown below.

Figure 4.1. Main Data Window

For more information see the ParaProf section in the reference guide.

4.3. Jumpshot

To use Argonne's Jumpshot (bundled with TAU), first merge and convert TAU traces to slog2 format:

```
% tau_treemerge.pl
% tau2slog2 tau.trc tau.edf -o tau.slog2
% jumpshot tau.slog2
```

Launching Jumpshot will bring up the main display window showing the entire trace, zoom in to see more detail.

Chapter 5. Quick Reference

```
tau run
   TAU's binary instrumentation tool
                  -tau_options=-optCompInst
tau_options=-optCompInst / tau_f90.sh -tau_options=-optCompInst
   Compiler wrappers (Compiler instrumentation)
tau_cc.sh / tau_cxx.sh / tau_f90.sh
   Compiler wrappers (PDT instrumentation)
TAU_MAKEFILE
   Set instrumentation definition file
TAU OPTIONS
   Set instrumentation options
dynamic phase
line=end_line_#
                    name='name'
                                   file='filename'
                                                          line=start line #
   Specify dynamic Phase
loops file='filename' routine='routine name'
   Instrument outer loops
memory file='filename' routine='routine name'
   Track memory
io file='filename' routine='routine name'
   Track IO
TAU_PROFILE / TAU_TRACE
   Enable profiling and/or tracing
PROFILEDIR / TRACEDIR
   Set profile/trace output directory
TAU_CALLPATH=1 / TAU_CALLPATH_DEPTH
   Enable Callpath profiling, set callpath depth
TAU THROTTLE=1 / TAU THROTTLE NUMCALLS / TAU THROTTLE PERCALL
   Enable event throttling, set number of call, percall (us) threshold
TAU METRICS
   List of PAPI metrics to profile
tau_treemerge.pl
   Merge traces to one file
tau2otf / tau2vtf / tau2slog2
```

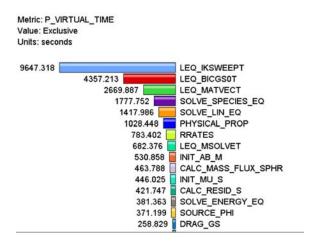
Trace conversion tools

Chapter 6. Some Common Application Scenario

6.1. Q. What routines account for the most time? How much?

A. Create a flat profile with wallclock time.

Figure 6.1. Flat Profile

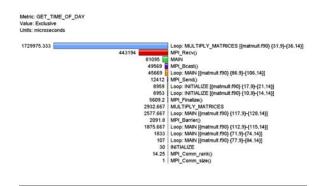


Here is how to generate a flat profile with MPI

6.2. Q. What loops account for the most time? How much?

A. Create a flat profile with wallclock time with loop instrumentation.

Figure 6.2. Flat Profile with Loops

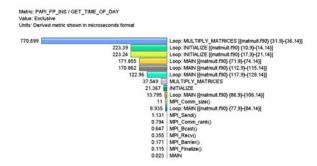


Here is how to instrument loops in an application

6.3. Q. What MFlops am I getting in all loops?

A. Create a flat profile with PAPI_FP_INS/OPS and time with loop instrumentation.

Figure 6.3. MFlops per loop

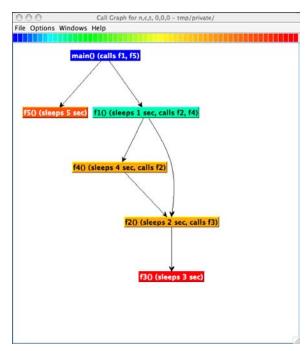


Here is how to generate a flat profile with FP operations

6.4. Q. Who calls MPI_Barrier() Where?

A. Create a callpath profile with given depth.





Here is how to generate a callpath profile with MPI

```
% setenv TAU_MAKEFILE
% /opt/apps/tau/tau-2.17.1/x86_64/lib/Makefile.tau-mpi-pdt
% set path=(/opt/apps/tau/tau-2.17.1/x86_64/bin $path)
% make F90=tau_f90.sh
(Or edit Makefile and change F90=tau_f90.sh)
```

6.5. Q. How do I instrument Python Code?

A. Create an python wrapper library.

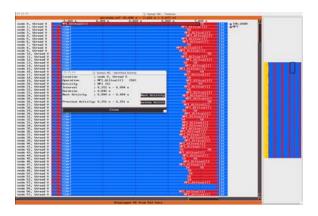
Here to instrument python code

```
% setenv TAU_MAKEFILE /opt/apps/tau/tau-2.17.1/x86_64/lib/Makefile.tau-icpc-python
% set path=(/opt/apps/tau/tau-2.17.1/x86_64/bin $path)
% setenv TAU OPTIONS `-optShared -optVerbose'
(Python needs shared object based TAU library)
% make F90=tau_f90.sh CXX=tau_cxx.sh CC=tau_cc.sh (build pyMPI w/TAU)
% cat wrapper.py
  import tau
  def OurMain():
      import App
  tau.run('OurMain()')
Uninstrumented:
% mpirun.lsf /pyMPI-2.4b4/bin/pyMPI ./App.py
Instrumented:
% setenv PYTHONPATH<taudir>/x86_64/<lib>/bindings-python-mpi-pdt-pgi
(same options string as TAU_MAKEFILE)
setenv LD_LIBRARY_PATH <taudir>/x86_64/lib/bindings-icpc-python-mpi-pdt-pgi\:$LD_L
% mpirun -np 4 <dir>/pyMPI-2.4b4-TAU/bin/pyMPI ./wrapper.py
(Instrumented pyMPI with wrapper.py)
```

6.6. Q. What happens in my code at a given time?

A. Create an event trace.

Figure 6.5. Tracing with Vampir



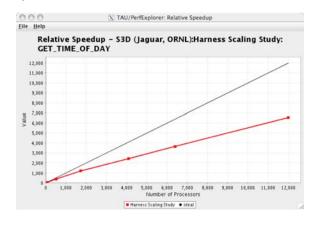
How to create a trace

```
% setenv TAU_MAKEFILE
% /opt/apps/tau/tau-2.17.1/x86_64/lib/Makefile.tau-mpi-pdt-pgi
% set path=(/opt/apps/tau/tau-2.17.1/x86_64/bin $path)
% make F90=tau_f90.sh
(Or edit Makefile and change F90=tau_f90.sh)
% setenv TAU_TRACE 1
% qsub run.job
% tau_treemerge.pl
(merges binary traces to create tau.trc and tau.edf files)
JUMPSHOT:
% tau2slog2 tau.trc tau.edf -o app.slog2
% jumpshot app.slog2
   OR
VAMPIR:
% tau2otf tau.trc tau.edf app.otf -n 4 -z
(4 streams, compressed output trace)
% vampir app.otf
(or vng client with vngd server).
```

6.7. Q. How does my application scale?

A. Examine profiles in PerfExplorer.

Figure 6.6. Scalability chart



How to examine a series of profiles in PerfExplorer

```
% setenv TAU_MAKEFILE /opt/apps/tau/tau-2.17.1/x86_64/lib/Makefile.tau-mpi-pdt
% set path=(/opt/apps/tau/tau-2.17.1/x86_64/bin $path)
% make F90=tau_f90.sh
(Or edit Makefile and change F90=tau_f90.sh)
% qsub runlp.job
% paraprof --pack 1p.ppk
% qsub run2p.job
% paraprof --pack 2p.ppk ...and so on.
On your client:
% perfdmf_configure
(Choose derby, blank user/password, yes to save password, defaults)
% perfexplorer_configure
(Yes to load schema, defaults)
% paraprof
(load each trial: Right click on trial ->Upload trial to DB
% perfexplorer
(Charts -> Speedup)
```

ParaProf - User's Manual

University of Oregon

ParaProf - User's Manual

by University of Oregon

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Chapter 1. Introduction

ParaProf is a portable, scalable performance analysis tool included with the TAU distribution.



Important

ParaProf requires Sun's Java 1.3 Runtime Environment for basic functionality. Java 1.4 is required for 3d visualization and image export. Additionally, OpenGL is required for 3d visualization.



Note

Most windows in ParaProf can export bitmap (png/jpg) and vector (svg/eps) images to disk (png/jpg) or print directly to a printer. This are available through the *File* menu.

1.1. Using ParaProf from the command line

ParaProf is a java program that is run from the supplied **paraprof** script (**paraprof.bat** for windows binary release).

```
% paraprof --help
Usage: paraprof [options] <files/directory>
Options:
  -f, --filetype <filetype>
                              Specify type of performance data,
                                options are: profiles (default), pprof,
                                dynaprof, mpip, gprof, psrun, hpm,
                                packed, cube, hpc, ompp
  -h, --help
                              Display this help message
                              Use `pprof` to compute derived data
  -i, --fixnames
                              Use the fixnames option for gprof
  -m, --monitor
                              Perform runtime monitoring of profile data
The following options will run only from the console (no GUI will launch):
  --pack <file>
                              Pack the data into packed (.ppk) format
                              Dump profile data to TAU profile format
  --dump
  -o, --oss
                              Print profile data in OSS style text output
  -s, --summary
                              Print only summary statistics
                                 (only applies to OSS output)
```

Notes:

For the TAU profiles type, you can specify either a specific set of profile files on the commandline, or you can specify a directory (by default the current directory). The specified directory will be searched for profile.*.*.* files, or, in the case of multiple counters, directories named MULTI_* containing profile data.

1.2. Supported Formats

ParaProf can load profile date from many sources. The types currently supported are:

- TAU Profiles Output from the TAU measurement library, these files generally take the form of profile.X.X.X, one for each node/context/thread combination. When multiple counters are used, each metric is located in a directory prefixed with "MULTI_". To launch ParaProf with all the metrics, simply launch it from the root of the MULTI_ directories.
- **pprof** Dump Output from TAU's **pprof** -d. Provided for backward compatibility only.
- **DynaProf** Output From DynaProf's wallclock and papi probes.
- mpiP Output from mpiP.
- **gprof** Output from gprof, see also the --fixnames option.
- **HPM Toolkit** Output from HPM Toolkit.
- ParaProf Packed Format Export format supported by PerfDMF/ParaProf. Typically .ppk.
- **Cube** Output from Kojak Expert tool for use with Cube.
- HPCToolkit XML data from hpcquick. Typically, the user runs hpcrun, then hpcquick on the resulting binary file.
- **ompP** CSV format from the ompP OpenMP Profiler (http://www.ompp-tool.com). The user must use OMPP_OUTFORMAT=CVS.

1.3. Command line options

In addition to specifying the profile format, the user can also specify the following options

- --fixnames Use the fixnames option for gprof. When C and Fortran code are mixed, the C routines
 have to be mapped to either .function or function_. Strip the leading period or trailing underscore, if
 it is there.
- --pack <file> Rather than load the data and launch the GUI, pack the data into the specified file.
- --dump Rather than load the data and launch the GUI, dump the data to TAU Profiles. This can be used to convert supported formats to TAU Profiles.
- --oss Outputs profile data in OSS Style. Example:

Thread: n,c,t 0,0,0									
excl.secs	excl.%	cum.%	PAPI_TOT_CYC	PAPI_FP_OPS	calls	function			
0.005	56.0%	56.0%	13475345	4194518	1	foo			
0.003	40.1%	96.1%	9682185	4205367		bar			
0	3.6%	99.7%	223173	17445	1	baz			
2.2E-05	0.3%	100.0%	14663	206		main			

• --summary - Output only summary information for OSS style output.

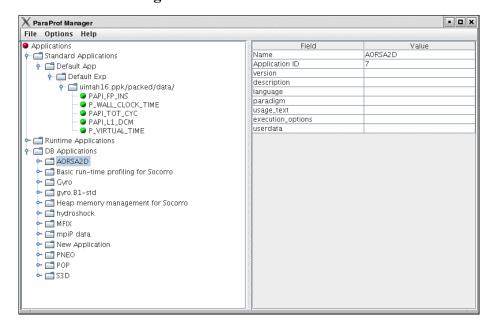
Chapter 2. Profile Data Management

ParaProf uses *PerfDMF* to manage profile data. This enables it to read the various profile formats as well as store and retrieve them from a database.

2.1. ParaProf Manager Window

Upon launching ParaProf, the user is greeted with the ParaProf Manager Window.

Figure 2.1. ParaProf Manager Window

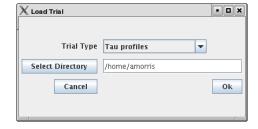


This window is used to manage profile data. The user can upload/download profile data, edit meta-data, launch visual displays, export data, derive new metrics, etc.

2.2. Loading Profiles

To load profile data, select File->Open, or right click on the Application's tree and select "Add Trial".

Figure 2.2. Loading Profile Data



Select the type of data from the "Trial Type" drop-down box. For TAU Profiles, select a directory, for other types, files.

2.3. Database Interaction

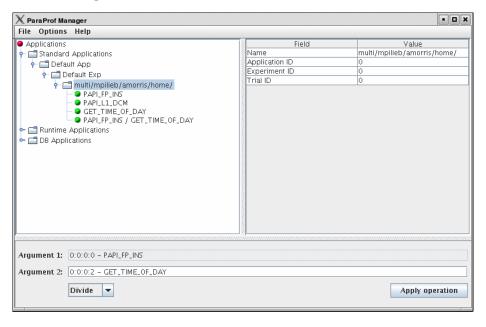
Database interaction is done through the tree view of the ParaProf Manager Window. Applications expand to Experiments, Experiments to Trials, and Trials are loaded directly into ParaProf just as if they were read off disk. Additionally, the meta-data associated with each element is show on the right, as in Figure 2.1, "ParaProf Manager Window". A trial can be exported by right clicking on it and selecting "Export as Packed Profile".

New trials can be uploaded to the database by either right-clicking on an entity in the database and selecting "Add Trial", or by right-clicking on an Application/Experiment/Trial hierarchy from the "Standard Applications" and selecting "Upload Application/Experiment/Trial to DB".

2.4. Creating Derived Metrics

ParaProf can created derived metrics using the *Derived Metric Panel*, available from the *Options* menu of the ParaProf Manager Window.

Figure 2.3. Creating Derived Metrics



In Figure 2.3, "Creating Derived Metrics", we have just divided Floating Point Instructions by Wall-clock time, creating FLOPS (Floating Point Operations per Second). The 2nd argument is a user editable text-box and can be filled in with scalar values by using the keyword 'val' (e.g. "val 1.5").

2.5. Main Data Window

Upon loading a profile, or double-clicking on a metric, the *Main Data Window* will be displayed.

Figure 2.4. Main Data Window

This window shows each thread as well as statistics as a combined bar graph. Each function is represented by a different color (though possibly cycled). From anywhere in ParaProf, you can right-click on objects representing threads or functions to launch displays associated with those objects. For example, in Figure 2.4, "Main Data Window", right click on the text n, c, t, 8, 0, 0 to launch thread based displays for node 8.

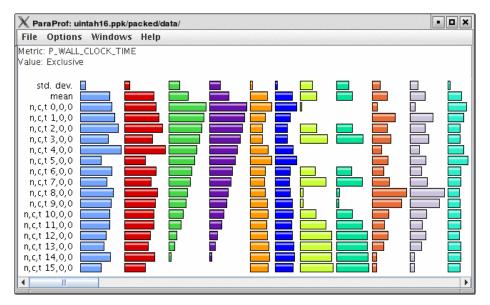


Figure 2.5. Unstacked Bars

You may also turn off the stacking of bars so that individual functions can be compared across threads in a global display.

Chapter 3. 3-D Visualization

ParaProf displays massive parallel profiles through the use of OpenGL hardware acceleration through the *3D Visualization* window. Each window is fully configurable with rotation, translation, and zooming capabilities. Rotation is accomplished by holding the left mouse button down and dragging the mouse. Translation is done likewise with the right mouse button. Zooming is done with the mousewheel and the + and - keyboard buttons.

3.1. Triangle Mesh Plot

X ParaProf Visualizer: Application 13, Experiment 23, Trial 58 - 0 × File Options Windows Help Triangle Mesh Bar Plot Scatter Plot Height Metric 14.538 ▼ Time • Exclusive Color Metric Exclusive Time MPI_Recv() Function 351:0:0 Thread Height value 14.37 seconds Color value Mesh Plot Axes ColorScale Render Plot Width Plot Denth Plot Height Transparency

Figure 3.1. Triangle Mesh Plot

This visualization method shows two metrics for all functions, all threads. The height represents one chosen metric, and the color, another. These are selected from the drop-down boxes on the right.

To pinpoint a specific value in the plot, move the *Function* and *Thread* sliders to cycle through the available functions/threads. The values for the two metrics, in this case for MPI_Recv() on Node 351, the value is 14.37 seconds.

3.2. 3-D Bar Plot

Figure 3.2. 3-D Mesh Plot

This visualization method is similar to the triangle mesh plot. It simply displays the data using 3d bars instead of a mesh. The controls works the same. Note that in Figure 3.2, "3-D Mesh Plot" the transparency option is selected, which changes the way in which the selection model operates.

3.3. 3-D Scatter Plot

- 0 × X ParaProf Visualizer: Application 13, Experiment 23, Trial 58. Triangle Mesh Bar Plot 0.221 Scatter Plot MPI_Send() Width ▼ Time ▼ Exclusive MPI_Recv() Depth Exclusive ▼ Time ▼ buts Height Exclusive ▼ Time ▼ exchange_3 Exclusive ▼ Time ▼

Figure 3.3. 3-D Scatter Plot

This visualization method plots the value of each thread along up to 4 axes (each a different function/metric). This view allows you to discern clustering of values and relationships between functions across threads.

ScatterPlot Axes ColorScale Render

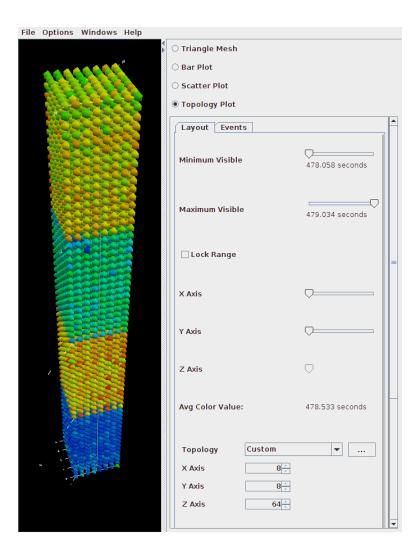
Point size Point detail

Select functions using the button for each dimension, then select a metric. A single function across 4 metrics could be used, for example.

3.4. 3-D Topology Plot

0.084

Figure 3.4. 3-D Topology Plot



In this visualization, you can either define the layout with a MESP topology definition file or you can fill a rectangular prism of user-defined volume with rank-points in order of rank. For more information, please see the etc/topology directory for additional details on MESP topology definitions.

If the loaded profile is a cube file or a profile from a BGB, then this visualizations groups the threads in two or three dimensional space using topology information supplied by the profile.

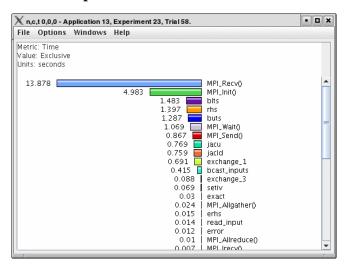
The right side bar provides options to manipulate the visualization. You can select different metrics or topologies. The sliders toward the top allow you to select the range of points to show.

Chapter 4. Thread Based Displays

ParaProf displays several windows that show data for one thread of execution. In addition to per thread values, the users may also select *mean* or *standard deviation* as the "thread" to display. In this mode, the mean or standard deviation of the values across the threads will be used as the value.

4.1. Thread Bar Graph

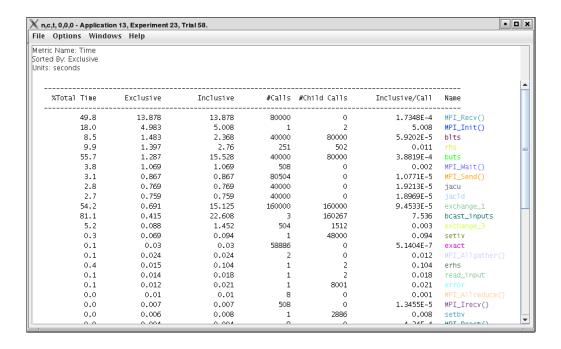
Figure 4.1. Thread Bar Graph



This display graphs each function on a particular thread for comparison. The metric, units, and sort order can be changed from the *Options* menu.

4.2. Thread Statistics Text Window

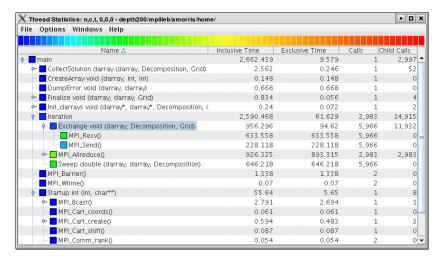
Figure 4.2. Thread Statistics Text Window



This display shows a **pprof** style text view of the data.

4.3. Thread Statistics Table

Figure 4.3. Thread Statistics Table, inclusive and exclusive



This display shows the callpath data in a table. Each callpath can be traced from root to leaf by opening each node in the tree view. A colorscale immediately draws attention to "hot spots", areas that contain highest values.

Figure 4.4. Thread Statistics Table

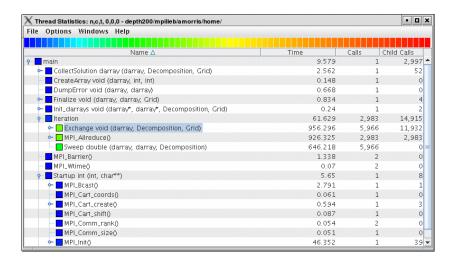
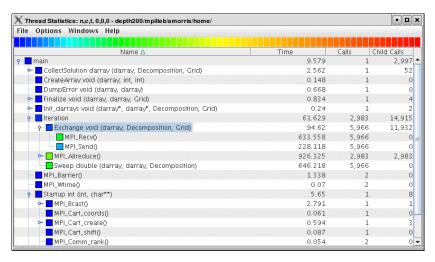


Figure 4.5. Thread Statistics Table

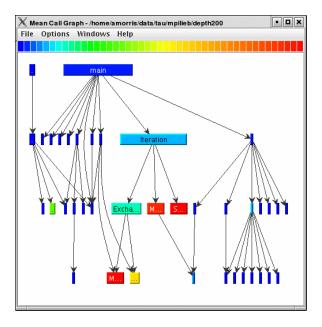


The display can be used in one of two ways, in "inclusive/exclusive" mode, both the inclusive and exclusive values are shown for each path, see Figure 4.3, "Thread Statistics Table, inclusive and exclusive" for an example.

When this option is off, the inclusive value for a node is show when it is closed, and the exclusive value is shown when it is open. This allows the user to more easily see where the time is spent since the total time for the application will always be represented in one column. See Figure 4.4, "Thread Statistics Table" and Figure 4.5, "Thread Statistics Table" for examples. This display also functions as a regular statistics table without callpath data. The data can be sorted by columns by clicking on the column heading. When multiple metrics are available, you can add and remove columns for the display using the menu.

4.4. Call Graph Window

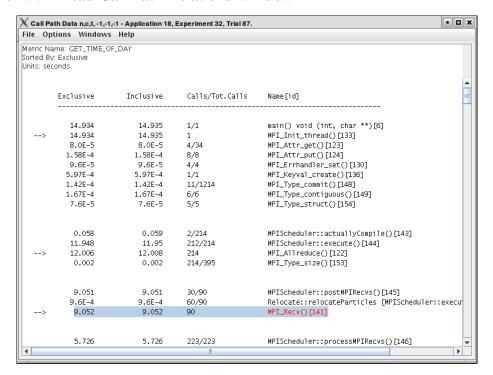
Figure 4.6. Call Graph Window



This display shows callpath data in a graph using two metrics, one determines the width, the other the color. The full name of the function as well as the two values (color and width) are displayed in a tooltip when hovering over a box. By clicking on a box, the actual ancestors and descendants for that function and their paths (arrows) will be highlighted with blue. This allows you to see which functions are called by which other functions since the interplay of multiple paths may obscure it.

4.5. Thread Call Path Relations Window

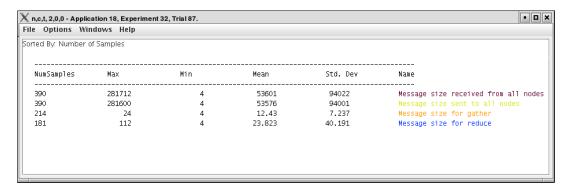
Figure 4.7. Thread Call Path Relations Window



This display shows callpath data in a **gprof** style view. Each function is shown with its immediate parents. For example, Figure 4.7, "Thread Call Path Relations Window" shows that MPI_Recv() is call from two places for a total of 9.052 seconds. Most of that time comes from the 30 calls when MPI_Recv() is called by MPIScheduler::postMPIRecvs(). The other 60 calls do not amount to much time.

4.6. User Event Statistics Window

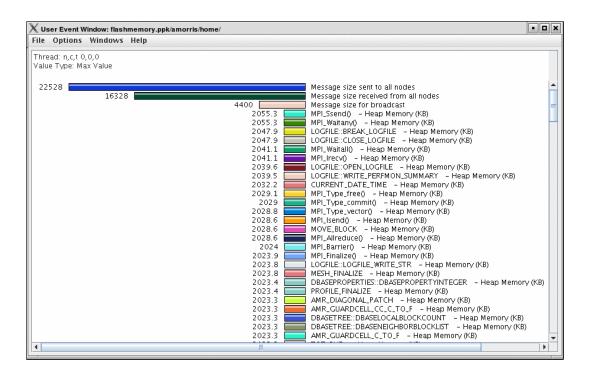
Figure 4.8. User Event Statistics Window



This display shows a **pprof** style text view of the user event data. Right clicking on a User Event will give you the option to open a Bar Graph for that particular User Event across all threads. See Section 8.1, "User Event Bar Graph"

4.7. User Event Thread Bar Chart

Figure 4.9. User Event Thread Bar Chart Window



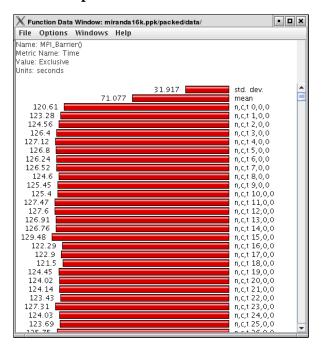
This display shows a particular thread's user defined event statistics as a bar chart. This is the same data from the Section 4.6, "User Event Statistics Window", in graphical form.

Chapter 5. Function Based Displays

ParaProf has two displays for showing a single function across all threads of execution. This chapter describes the Function Bar Graph Window and the Function Histogram Window.

5.1. Function Bar Graph

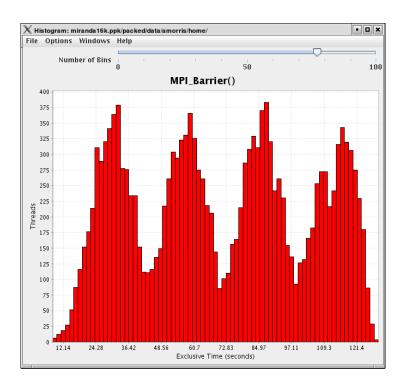
Figure 5.1. Function Bar Graph



This display graphs the values that the particular function had for each thread along with the mean and standard deviation across the threads. You may also change the units and metric displayed from the *Options* menu.

5.2. Function Histogram

Figure 5.2. Function Histogram



This display shows a histogram of each thread's value for the given function. Hover the mouse over a given bar to see the range minimum and maximum and how many threads fell into that range. You may also change the units and metric displayed from the *Options* menu.

You may also dynamically change how many bins are used (1-100) in the histogram. This option is available from the *Options* menu. Changing the number of bins can dramatically change the shape of the histogram, play around with it to get a feel for the true distribution of the data.

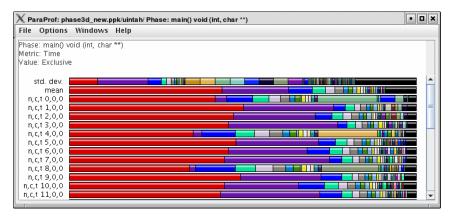
Chapter 6. Phase Based Displays

When a profile contains phase data, ParaProf will automatically run in phase mode. Most displays will show data for a particular phase. This phase will be displayed in teh top left corner in the meta data panel.

6.1. Using Phase Based Displays

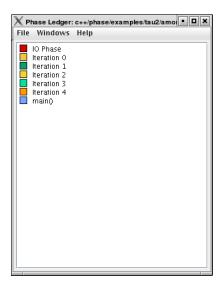
The initial window will default to top level phase, usually main

Figure 6.1. Initial Phase Display



To access other phases, either right click on the phase and select, "Open Profile for this Phase", or go to the *Phase Ledger* and select it there.

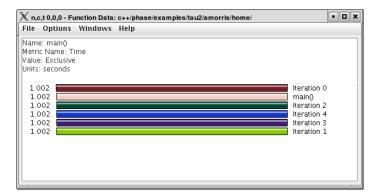
Figure 6.2. Phase Ledger



ParaProf can also display a particular function's value across all of the phases. To do so, right click on a

function and select, "Show Function Data over Phases".

Figure 6.3. Function Data over Phases



Because Phase information is implemented as callpaths, many of the callpath displays will show phase data as well. For example, the Call Path Text Window is useful for showing how functions behave across phases.

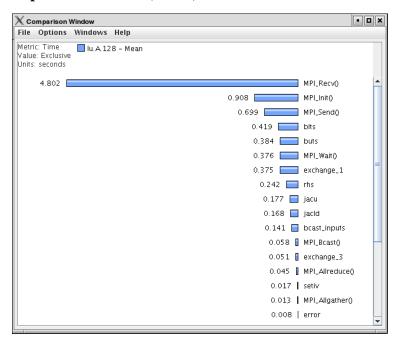
Chapter 7. Comparative Analysis

ParaProf can perform cross-thread and cross-trial analysis. In this way, you can compare two or more trials and/or threads in a single display.

7.1. Using Comparitive Analysis

Comparative analysis in ParaProf is based on individual threads of execution. There is a maximum of one Comparison window for a given ParaProf session. To add threads to the window, right click on them and select "Add Thread to Comparison Window". The Comparison Window will pop up with the thread selected. Note that "mean" and "std. dev." are considered threads for this any most other purposes.

Figure 7.1. Comparison Window (initial)



Add additional threads, from any trial, by the same means.

Figure 7.2. Comparison Window (2 trials)

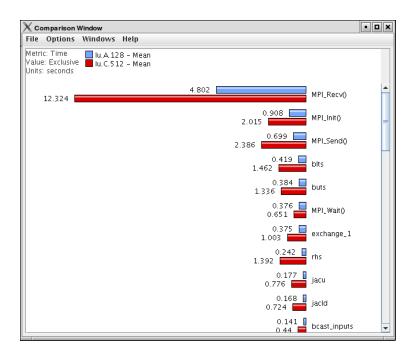
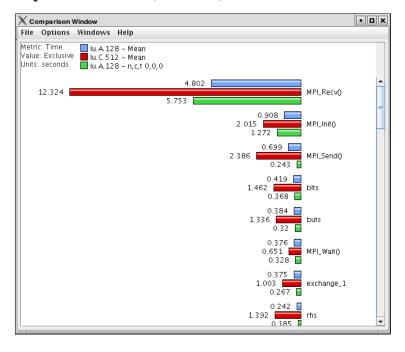


Figure 7.3. Comparison Window (3 threads)

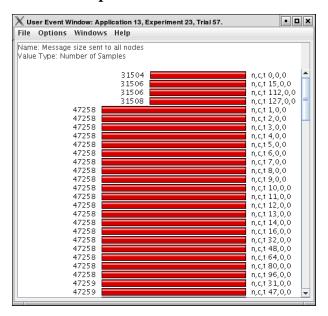


Chapter 8. Miscellaneous Displays

8.1. User Event Bar Graph

In addition to displaying the text statistics for User Defined Events, ParaProf can also graph a particular User Event across all threads.

Figure 8.1. User Event Bar Graph



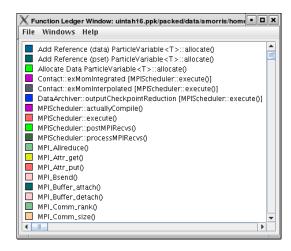
This display graphs the value that the particular user event had for each thread.

8.2. Ledgers

ParaProf has three ledgers that show the functions, groups, and user events.

8.2.1. Function Ledger

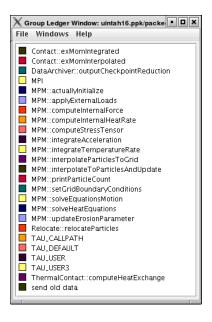
Figure 8.2. Function Ledger



The function ledger shows each function along with its current color. As with other displays showing functions, you may right-click on a function to launch other function-specific displays.

8.2.2. Group Ledger

Figure 8.3. Group Ledger



The group ledger shows each group along with its current color. This ledger is especially important because it gives you the ability to mask all of the other displays based on group membership. For example, you can right-click on the MPI group and select "Show This Group Only" and all of the windows will now mask to only those functions which are members of the MPI group. You may also mask by the inverse by selecting "Show All Groups Except This One" to mask out a particular group.

8.2.3. User Event Ledger

Figure 8.4. User Event Ledger

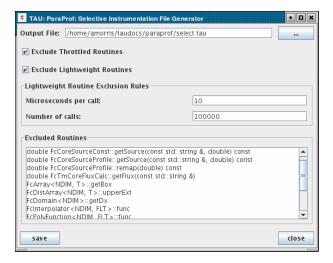


The user event ledger shows each user event along with its current color.

8.3. Selective Instrumentation File Generator

ParaProf can also help you refine your program performance by excluding some functions from instrumentation. You can select rules to determine which function get excluded; both rules must be true for a given function to be excluded. Below each function that will be excluded based on these rules are listed.

Figure 8.5. Selective Instrumentation Dialog





Note

Only the functions profilied in ParaProf can be excluded. If you had previously setup selective instrumentation for this application the functions that where previously excluded will not longer be excluded.

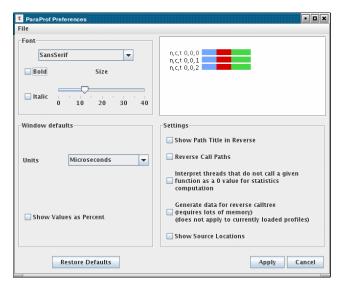
Chapter 9. Preferences

Preferences are modified from the ParaProf Preferences Window, launched from the File menu. Preferences are saved between sessions in the .ParaProf/ParaProf.prefs

9.1. Preferences Window

In addition to displaying the text statistics for User Defined Events, ParaProf can also graph a particular User Event across all threads.

Figure 9.1. ParaProf Preferences Window



The preferences window allows the user to modify the behavior and display style of ParaProf's windows. The font size affects bar height, a sample display is shown in the upper-right.

The Window defaults section will determine the initial settings for new windows. You may change the initial units selection and whether you want values displayed as percentages or as raw values.

The Settings section controls the following

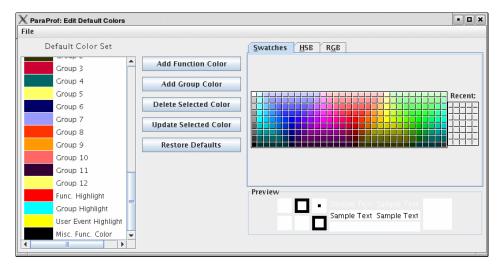
- Show Path Title in Reverse Path title will normally be shown in normal order (/home/amorris/data/etc). They can be reverse using this option (etc/data/amorris/home). This only affects loaded trials and the titlebars of new windows.
- Reverse Call Paths This option will immediately change the display of all callpath functions between Root => Leaf and Leaf <= Root.
- Statistics Computation Turning this option on causes the mean computation to take the sum of
 value for a function across all threads and divide it by the total number of threads. With this option
 off the sum will only be divided by the number of threads that actively participated in the sum. This
 way the user can control whether or not threads which do not call a particular function are consider
 as a 0 in the computation of statistics.
- · Generate Reverse Calltree Data This option will enable the generation of reverse callpath data ne-

cessary for the reverse callpath option of the statistics tree-table window.

 Show Source Locations - This option will enable the display of source code locations in event names.

9.2. Default Colors

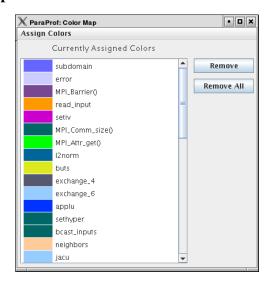
Figure 9.2. Edit Default Colors



The default color editor changes how colors are distributed to functions whose color has not been specifically assigned. It is accessible from the File menu of the Preferences Window.

9.3. Color Map

Figure 9.3. Color Map



The color map shows specifically assigned colors. These values are used across all trials loaded so that the user can identify a particular function across multiple trials. In order to map an entire trial's function set, Select "Assign Defaults from ->" and select a loaded trial.

Individual functions can be assigned a particular color by clicking on them in any of the other ParaProf Windows.

PerfExplorer - User's Manual

University of Oregon

PerfExplorer - User's Manual by University of Oregon

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Chapter 1. Introduction

PerfExplorer is a framework for parallel performance data mining and knowledge discovery. The framework architecture enables the development and integration of data mining operations that will be applied to large-scale parallel performance profiles.

The overall goal of the PerfExplorer project is to create a software to integrate sophisticated data mining techniques in the analysis of large-scale parallel performance data.

PerfExplorer supports clustering, summarization, association, regression, and correlation. Cluster analysis is the process of organizing data points into logically similar groupings, called clusters. Summarization is the process of describing the similarities within, and dissimilarities between, the discovered clusters. Association is the process of finding relationships in the data. One such method of association is regression analysis, the process of finding independent and dependent correlated variables in the data. In addition, comparative analysis extends these operations to compare results from different experiments, for instance, as part of a parametric study.

In addition to the data mining operations available, the user may optionally choose to perform comparative analysis. The types of charts available include time-steps per second, relative efficiency and speedup of the entire application, relative efficiency and speedup for all events, relative efficiency and speedup for all phases and runtime breakdown of the application by event or by phase. In addition, when the events are grouped together, such as in the case of communication routines, yet another chart shows the percentage of total runtime spent in that group of events. These analyses can be conducted across different combinations of parallel profiles and across phases within an execution.

Chapter 2. Installation and Configuration

PerfExplorer uses PerfDMF databases so if you have not already you will need to install PerfDMF, see ???. To configure PerfExplorer move to the tools/src/PerfExplorer/ directory in you TAU distribution. Type:

%>./configure

If you haven't already done so for other TAU tools, add [path to tau]/tau2/apple/bin to your path.

The following command-line options are available to configure:

2.1. Available configuration options

-engine=<analysis engine>

Specifies the data-mining engine to use. The supported options include weka and R.

• -rroot=<directory>

Specifies the directory where R is installed. Specifically, it should be the directory where the bin, include, lib, library and share directories are located.

• -objectport=<available network port>

Specifies the port that the PerfExplorer server should use, when running PerfExplorer in client-server mode. Select an available network port, and make sure that other appropriate network configurations are made (firewalls, etc.). The default port is 9999.

• -registryport=<available network port>

Specifies the port that the rmiregistry should use, when ruining PerfExplorer in client-server mode. Select an available network port, and make sure that other appropriate network configurations are made (firewalls, etc.). The default port is 1099.

-server=<server name>

Specifies the fully qualified domain name of the server where PerfExplorer is run, when running PerfExplorer in client-server mode.

Chapter 3. Running PerfExplorer

To run PerfExplorer type:

%>perfexplorer

When PerfExplorer loads you will see on the left window all the experiments that where loaded into PerfDMF. You can select which performance data you are interested by navigating the tree structure. PerfExplorer will allow you to run analysis operations on these experiments. Also the cluster analysis results are visible on the right side of the window. Various types of comparative analysis are available from the drop down menu selected.

To run an analysis operation, first select the metric of interest form the experiments on the left. Then perform the operation by selecting it from the Analysis menu. If you would like you can set the clustering method, dimension reduction, normalization method and the number of clusters from the same menu.

The options under the Charts menu provide analysis over one or more applications, experiments, views or trials. To view these charts first choose a metric of interest by selecting a trial form the tree on the left. Then optionally choose the Set Metric of Interest or Set Event of Interest form the Charts menu (if you don't, and you need to, you will be prompted). Now you can view a chart by selecting it from the Charts menu.

Chapter 4. Cluster Analysis

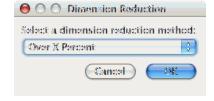
Cluster analysis is a valuable tool for reducing large parallel profiles down to representative groups for investigation. Currently, there are two types of clustering analysis implemented in PerfExplorer. Both hierarchical and k-means analysis are used to group parallel profiles into common clusters, and then the clusters are summarized. Initially, we used similarity measures computed on a single parallel profile as input to the clustering algorithms, although other forms of input are possible. Here, the performance data is organized into multi-dimensional vectors for analysis. Each vector represents one parallel thread (or process) of execution in the profile. Each dimension in the vector represents an event that was profiled in the application. Events can be any sub-region of code, including libraries, functions, loops, basic blocks or even individual lines of code. In simple clustering examples, each vector represents only one metric of measurement. For our purposes, some dissimilarity value, such as Euclidean or Manhattan distance, is computed on the vectors. As discussed later, we have tested hierarchical and \$k\$-means cluster analysis in PerfExplorer on profiles with over 32K threads of execution with few difficulties.

4.1. Dimension Reduction

Often, many hundreds of events are instrumented when profile data is collected. Clustering works best with dimensions less than 10, so dimension reduction is often necessary to get meaningful results. Currently, there is only one type of dimension reduction available in PerfExplorer. To reduce dimensions, the user specifies a minimum exclusive percentage for an event to be considered "significant".

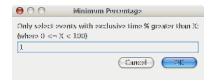
To reduce dimensions, select the "Select Dimension Reduction" item under the "Analysis" main menu bar item. The following dialog will appear:

Figure 4.1. Selecting a dimension reduction method



Select "Over X Percent". The following dialog will appear:

Figure 4.2. Entering a minimum threshold for exclusive percentage



Enter a value, for example "1".

4.2. Max Number of Clusters

By default, PerfExplorer will attempt k-means clustering with values of k from 2 to 10. To change the maximum number of clusters, select the "Set Maximum Number of Clusters" item under the "Analysis"

main menu item. The following dialog will appear:

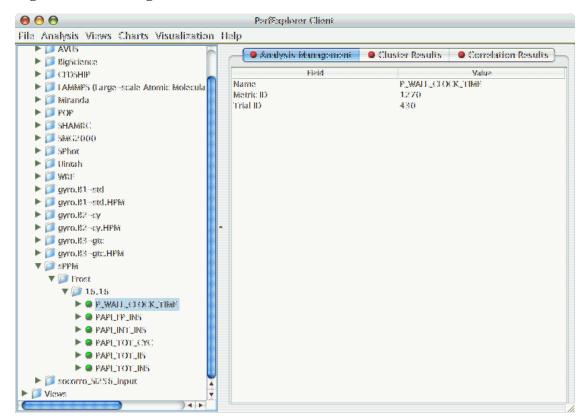
Figure 4.3. Entering a maximum number of clusters



4.3. Performing Cluster Analysis

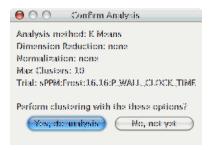
To perform cluster analysis, you first need to select a metric. To select a metric, navigate through the tree of applications, experiments and trials, and expand the trial of interest, showing the available metrics, as shown in the figure below:

Figure 4.4. Selecting a Metric to Cluster



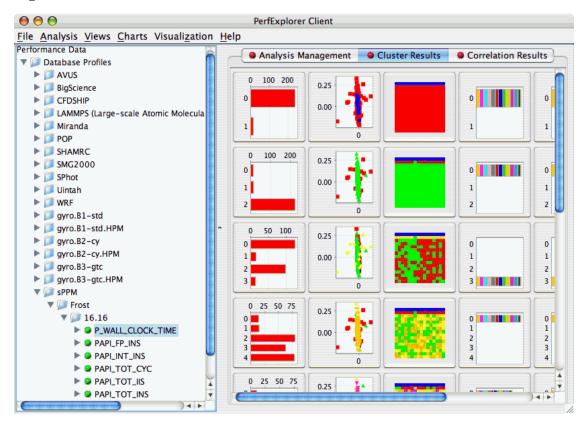
After selecting the metric of interest, select the "Do Clustering" item under the "Analysis" main menu bar item. The following dialog will appear:

Figure 4.5. Confirm Clustering Options



After confirming the clustering, the clustering will begin. When the clustering results are available, you can view them in the "Cluster Results" tab.

Figure 4.6. Cluster Results



There are a number of images in the "Cluster Results" window. From left to right, the windows indicate the cluster membership histogram, a PCA scatterplot showing the cluster memberships, a virtual topology of the parallel machine, the minimum values for each event in each cluster, the average values for each event in each cluster, and the maximum values for each event in each cluster. Clicking on a thumbnail image in the main window will bring up the images, as shown below:

Figure 4.7. Cluster Membership Histogram

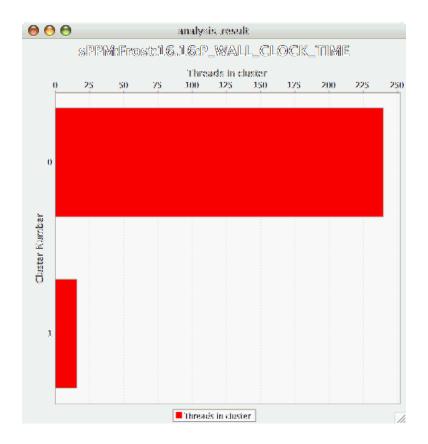


Figure 4.8. Cluster Membership Scatterplot

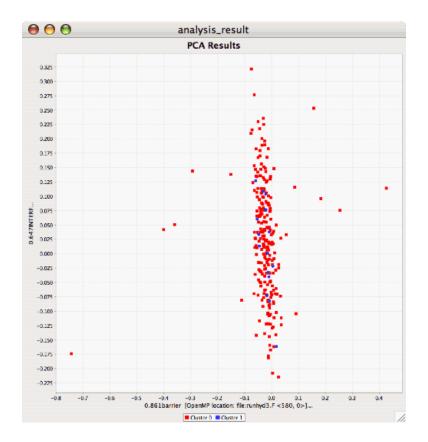


Figure 4.9. Cluster Virtual Topology

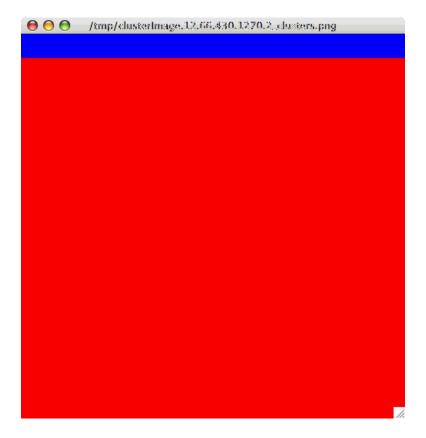
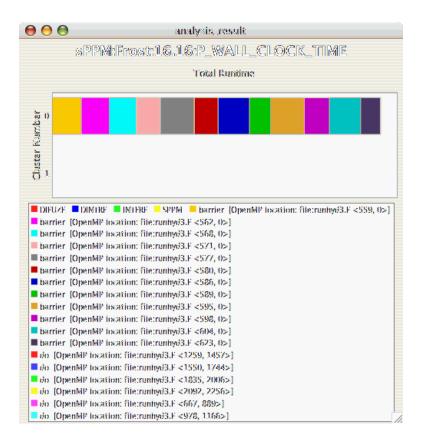


Figure 4.10. Cluster Average Behavior



Chapter 5. Correlation Analysis

Correlation analysis in PerfExplorer is used to explore relationships between events in a profile. Each event is pairwise plotted with the other events, and a correlation coefficient is calcuated for the relationship. When the events are highly positively correlated (coefficient of close to 1.0) or highly negatively correlated (coefficient close to -1.0), then the relationships will show up as linear groupings in the results. Clusters may also be apparent.

5.1. Dimension Reduction

Often, many hundreds of events are instrumented when profile data is collected. Clustering works best with dimensions less than 10, so dimension reduction is often necessary to get meaningful results. Currently, there is only one type of dimension reduction available in PerfExplorer. To reduce dimensions, the user specifies a minimum exclusive percentage for an event to be considered "significant".

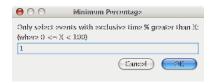
To reduce dimensions, select the "Select Dimension Reduction" item under the "Analysis" main menu bar item. The following dialog will appear:

Figure 5.1. Selecting a dimension reduction method



Select "Over X Percent". The following dialog will appear:

Figure 5.2. Entering a minimum threshold for exclusive percentage

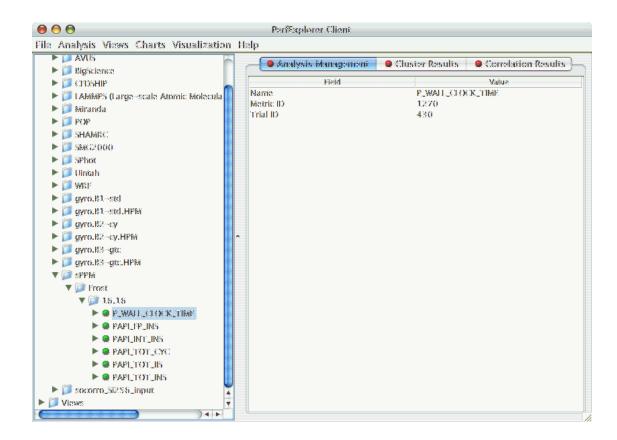


Enter a value, for example "1".

5.2. Performing Correlation Analysis

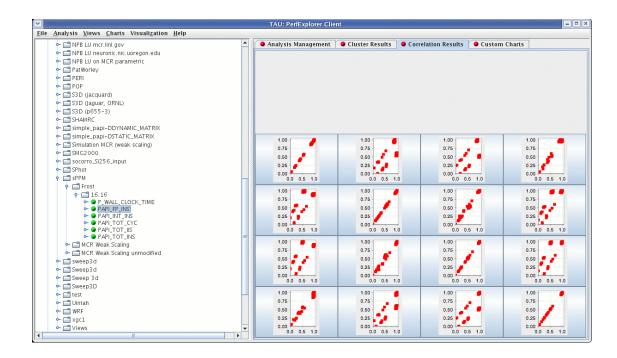
To perform correlation analysis, you first need to select a metric. To select a metric, navigate through the tree of applications, experiments and trials, and expand the trial of interest, showing the available metrics, as shown in the figure below:

Figure 5.3. Selecting a Metric to Cluster



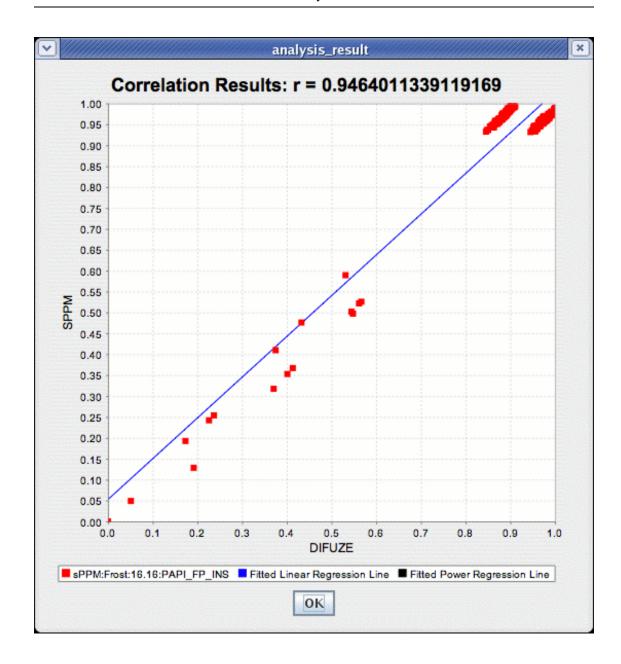
After selecting the metric of interest, select the "Do Correlation Analysis" item under the "Analysis" main menu bar item. A confirmation dialog will appear, and you can either confirm the correlation request or cancel it. After confirming the correlation, the analysis will begin. When the analysis results are available, you can view them in the "Correlation Results" tab.

Figure 5.4. Correlation Results



There are a number of images in the "Correlation Results" window. Each thumbnail represents a pairwise correlation plot of two events. Clicking on a thumbnail image in the main window will bring up the images, as shown below:

Figure 5.5. Correlation Example



Chapter 6. Charts

6.1. Setting Parameters

There are a few parameters which need to be set when doing comparisons between trials in the database. If any necessary setting is not configured before requesting a chart, you will be prompted to set the value. The following settings may be necessary for the various charts available:

6.1.1. Group of Interest

TAU events are often associated with common groups, such as "MPI", "TRANSPOSE", etc. This value is used for showing what fraction of runtime that this group of events contributed to the total runtime.

Figure 6.1. Setting Group of Interest



6.1.2. Metric of Interest

Profiles may contain many metrics gathered for a single trial. This selects which of the available metrics the user is interested in.

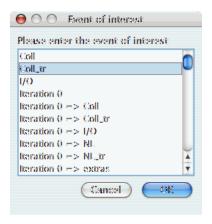
Figure 6.2. Setting Metric of Interest



6.1.3. Event of Interest

Some charts examine events in isolation. This setting configures which event to examine.

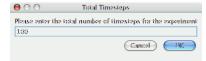
Figure 6.3. Setting Event of Interest



6.1.4. Total Number of Timesteps

One chart, the "Timesteps per second" chart, will calculate the number of timesteps completed per second. This setting configures that value.

Figure 6.4. Setting Timesteps

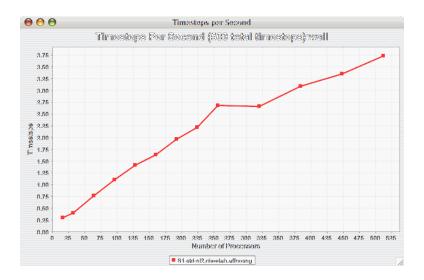


6.2. Standard Chart Types

6.2.1. Timesteps Per Second

The Timesteps Per Second chart shows how an application scales as it relates to time-to-solution. If the timesteps are not already set, you will be prompted to enter the total number of timesteps in the trial (see Section 6.1.4, "Total Number of Timesteps"). If there is more than one metric to choose from, you may be prompted to select the metric of interest (see Section 6.1.2, "Metric of Interest"). To request this chart, select one or more experiments or one view, and select this chart item under the "Charts" main menu item.

Figure 6.5. Timesteps per Second



6.2.2. Relative Efficiency

The Relative Efficiency chart shows how an application scales with respect to relative efficiency. That is, as the number of processors increases by a factor, the time to solution is expected to decrease by the same factor (with ideal scaling). The fraction between the expected scaling and the actual scaling is the relative efficiency. If there is more than one metric to choose from, you may be prompted to select the metric of interest (see Section 6.1.2, "Metric of Interest"). To request this chart, select one experiment or view, and select this chart item under the "Charts" main menu item.

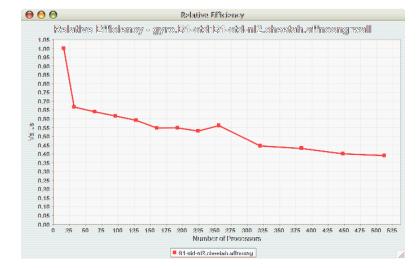


Figure 6.6. Relative Efficiency

6.2.3. Relative Efficiency by Event

The Relative Efficiency By Event chart shows how each event in an application scales with respect to relative efficiency. That is, as the number of processors increases by a factor, the time to solution is expected to decrease by the same factor (with ideal scaling). The fraction between the expected scaling and the actual scaling is the relative efficiency. If there is more than one metric to choose from, you may be prompted to select the metric of interest (see Section 6.1.2, "Metric of Interest"). To request this chart,

select one or more experiments or one view, and select this chart item under the "Charts" main menu item.

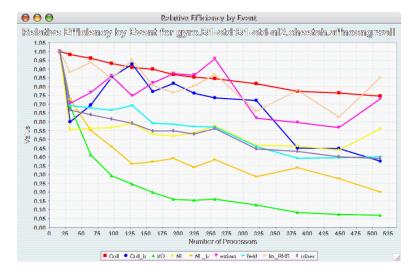


Figure 6.7. Relative Efficiency by Event

6.2.4. Relative Efficiency for One Event

The Relative Efficiency for One Event chart shows how one event from an application scales with respect to relative efficiency. That is, as the number of processors increases by a factor, the time to solution is expected to decrease by the same factor (with ideal scaling). The fraction between the expected scaling and the actual scaling is the relative efficiency. If there is more than one event to choose from, and you have not yet selected an event of interest, you may be prompted to select the event of interest (see Section 6.1.3, "Event of Interest"). If there is more than one metric to choose from, you may be prompted to select the metric of interest (see Section 6.1.2, "Metric of Interest"). To request this chart, select one or more experiments or one view, and select this chart item under the "Charts" main menu item.



Figure 6.8. Relative Efficiency one Event

6.2.5. Relative Speedup

The Relative Speedup chart shows how an application scales with respect to relative speedup. That is, as the number of processors increases by a factor, the speedup is expected to increase by the same factor (with ideal scaling). The ideal speedup is charted, along with the actual speedup for the application. If there is more than one metric to choose from, you may be prompted to select the metric of interest (see Section 6.1.2, "Metric of Interest"). To request this chart, select one or more experiments or one view, and select this chart item under the "Charts" main menu item.

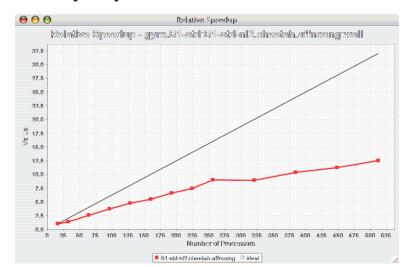
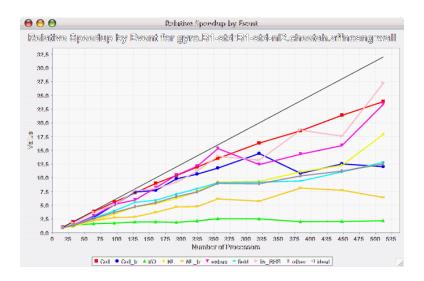


Figure 6.9. Relative Speedup

6.2.6. Relative Speedup by Event

The Relative Speedup By Event chart shows how the events in an application scale with respect to relative speedup. That is, as the number of processors increases by a factor, the speedup is expected to increase by the same factor (with ideal scaling). The ideal speedup is charted, along with the actual speedup for the application. If there is more than one metric to choose from, you may be prompted to select the metric of interest (see Section 6.1.2, "Metric of Interest"). To request this chart, select one experiment or view, and select this chart item under the "Charts" main menu item.

Figure 6.10. Relative Speedup by Event



6.2.7. Relative Speedup for One Event

The Relative Speedup for One Event chart shows how one event in an application scales with respect to relative speedup. That is, as the number of processors increases by a factor, the speedup is expected to increase by the same factor (with ideal scaling). The ideal speedup is charted, along with the actual speedup for the application. If there is more than one event to choose from, and you have not yet selected an event of interest, you may be prompted to select the event of interest (see Section 6.1.3, "Event of Interest"). If there is more than one metric to choose from, you may be prompted to select the metric of interest (see Section 6.1.2, "Metric of Interest"). To request this chart, select one or more experiments or one view, and select this chart item under the "Charts" main menu item.

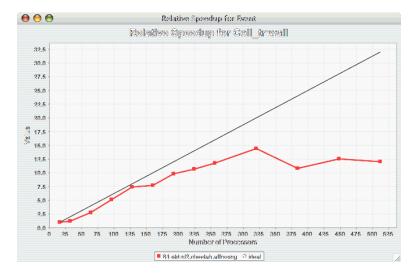


Figure 6.11. Relative Speedup one Event

6.2.8. Group % of Total Runtime

The Group % of Total Runtime chart shows how the fraction of the total runtime for one group of events changes as the number of processors increases. If there is more than one group to choose from, and you have not yet selected a group of interest, you may be prompted to select the group of interest (see Sec-

tion 6.1.1, "Group of Interest"). If there is more than one metric to choose from, you may be prompted to select the metric of interest (see Section 6.1.2, "Metric of Interest"). To request this chart, select one or more experiments or one view, and select this chart item under the "Charts" main menu item.

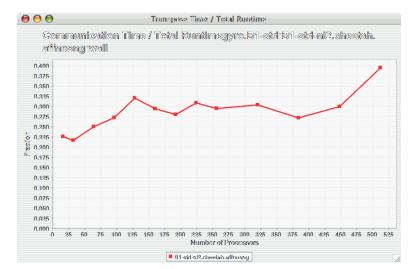


Figure 6.12. Group % of Total Runtime

6.2.9. Runtime Breakdown

The Runtime Breakdown chart shows the fraction of the total runtime for all events in the application, and how the fraction changes as the number of processors increases. If there is more than one metric to choose from, you may be prompted to select the metric of interest (see Section 6.1.2, "Metric of Interest"). To request this chart, select one experiment or view, and select this chart item under the "Charts" main menu item.

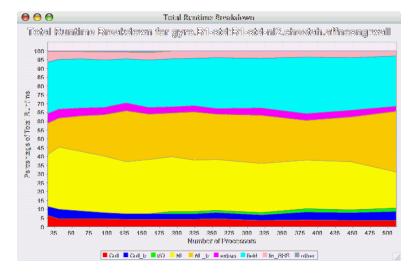


Figure 6.13. Runtime Breakdown

6.3. Phase Chart Types

TAU now provides the ability to break down profiles with respect to phases of execution. One such application would be to collect separate statistics for each timestep, or group of timesteps. In order to visualize the variance between the phases of execution, a number of phase-based charts are available.

6.3.1. Relative Efficiency per Phase

The Relative Efficiency Per Phase chart shows the relative efficiency for each phase, as the number of processors increases. If there is more than one metric to choose from, you may be prompted to select the metric of interest (see Section 6.1.2, "Metric of Interest"). To request this chart, select one experiment or view, and select this chart item under the "Charts" main menu item.

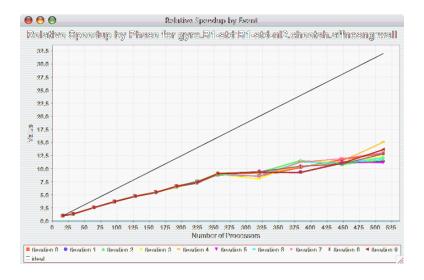


Figure 6.14. Relative Efficiency per Phase

6.3.2. Relative Speedup per Phase

The Relative Speedup Per Phase chart shows the relative speedup for each phase, as the number of processors increases. If there is more than one metric to choose from, you may be prompted to select the metric of interest (see Section 6.1.2, "Metric of Interest"). To request this chart, select one experiment or view, and select this chart item under the "Charts" main menu item.

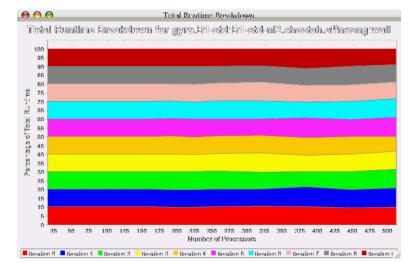
Figure 6.15. Relative Speedup per Phase



6.3.3. Phase Fraction of Total Runtime

The Phase Fraction of Total Runtime chart shows the breakdown of the execution by phases, and shows how that breakdown changes as the number of processors increases. If there is more than one metric to choose from, you may be prompted to select the metric of interest (see Section 6.1.2, "Metric of Interest"). To request this chart, select one experiment or view, and select this chart item under the "Charts" main menu item.

Figure 6.16. Phase Fraction of Total Runtime



Chapter 7. Custom Charts

In addition to the default charts available in the charts menu, there are is a custom chart interface. To access the interface, select the "Custom Charts" tab on in the results pane of the main window, as shown:

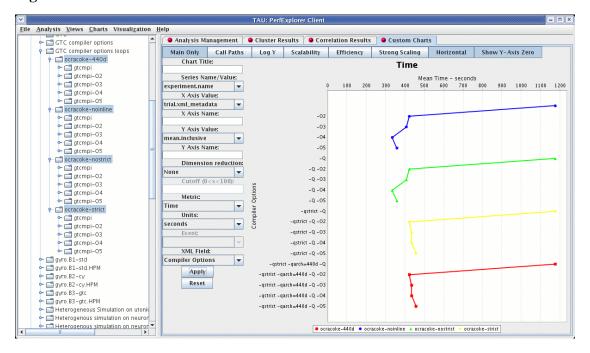


Figure 7.1. The Custom Charts Interface

There are a number of controls for the cusotom charts. They are:

- Main Only When selected, only the main event (the event with the highest inclusive value) will be selected. When deselected, the "Events" control (see below) is activated, and one or all events can be selected.
- Call Paths When selected, callpath events will be available in the "Events" control (see below).
- Log Y When selected, the Y axis will be the log of the true value.
- Scalability When selected, the chart will be interpreted as a speedup chart. The trial with the fewest number of threads of execution will be considered the baseline trial.
- Efficiency When selected, the chart will be interpreted as a relative efficiency chart. The trial with
 the fewest number of threads of execution will be considered the baseline trial.
- Strong Scaling When deselected, the speedup or efficiency chart will be interpreted as a strong
 scaling study (the workload is the same for all trials). When selected, the button will change to
 "Weak Scaling", and the chart will be interpreted as a weak scaling study (the workload is proportional to the total number of threads in each trial).
- Horizontal when selected, the chart X and Y axes will be swapped.
- Show Y-Axis Zero when selected, the chart will include the value 0. When deselected, the chart

will only show the relevant values for all data points.

- Chart Title value to use for the chart title
- Series Name/Value the field to be used to group the data points as a series.
- X Axis Value the field to use as the X axis value.
- X Axis Name the name to put in the chart for the value along the X axis.
- Y Axis Value the field to use as the Y axis value
- Y Axis Name the name to put in the chart for the value along the X axis.
- Dimension Reduction whether or not to use dimension reduction. This is only applicable when "Main Only" is disabled.
- Cutoff when the "Dimension Reduction" is enabled, the cutoff value for selecting "All Events".
- Metric The metric of interest for the Y axis.
- Units The unit to be selected for the Y axis.
- Event The event of interest, or "All Events".
- XML Field When the X or Y axis is selected to be an XML field, this is the field of interest.
- · Apply build the chart.
- Reset restore the options back to the default values.

When the chart is generated, it can be saved as a vector image by selecting "File -> Save As Vector Image". The chart can also be saved as a PNG by right clicking on the chart, and selecting "Save As...".

Chapter 8. Visualization

Under the "Visualization" main menu item, there are five types of raw data visualization. The five items are "3D Visualization", "Data Summary", "Create Boxchart", "Create Histogram" and "Create Normal Probability Chart". For the Boxchart, Histogram and Normal Probability Charts, you can either select one metric in the trial (which selects all events by default), or expand the metric and select events of interest

8.1. 3D Visualization

When the "3D Visualization" is requested, PerfExplorer examines the data to try to determine the most interesting events in the trial. That is, for the selected metric in the selected trial, the database will calculate the weighted relative variance for each event across all threads of execution, in order to find the top four "significant" events. These events are selected by calculating: stddev(exclusive) / (max(exclusive) - min(exclusive)) * exclusive_percentage. After selecting the top four events, they are graphed in an OpenGL window.

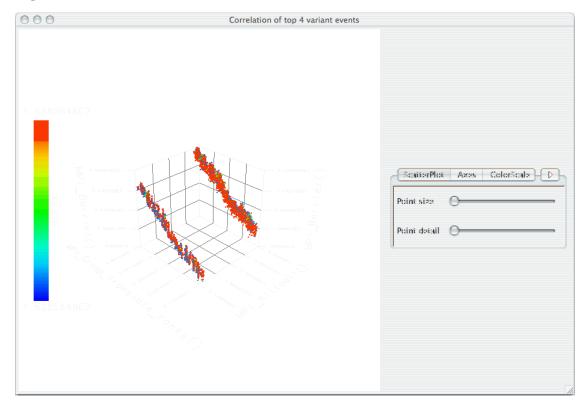
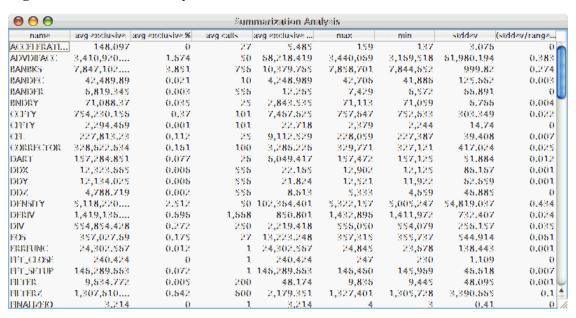


Figure 8.1. 3D Visualization of multivariate data

8.2. Data Summary

In order to see a summary of the performance data in the database, select the "Show Data Summary" item under the "Visualization" main menu item.

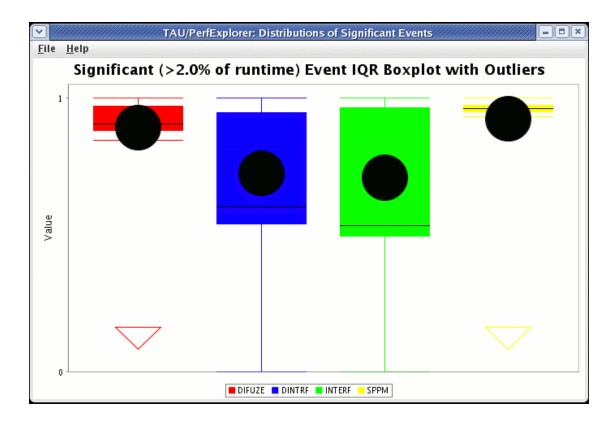
Figure 8.2. Data Summary Window



8.3. Creating a Boxchart

In order to see a boxchart summary of the performance data in the database, select the "Create Boxchart" item under the "Visualization" main menu item.

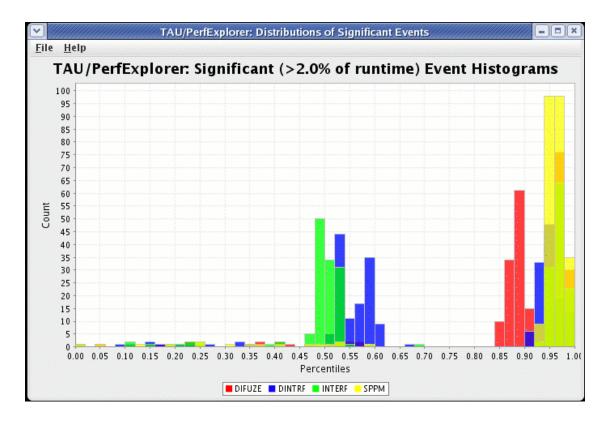
Figure 8.3. Boxchart



8.4. Creating a Histogram

In order to see a histogram summary of the performance data in the database, select the "Create Histogram" item under the "Visualization" main menu item.

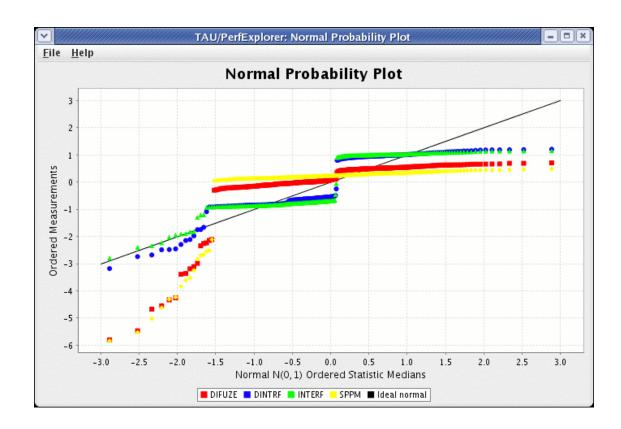
Figure 8.4. Histogram



8.5. Creating a Normal Probability Chart

In order to see a normal probability summary of the performance data in the database, select the "Create NormalProbability" item under the "Visualization" main menu item.

Figure 8.5. Normal Probability



Chapter 9. Views

Often times, data is loaded into the database with multiple parametric cross-sections. For example, the charts available in PerfExplorer are primarily designed for scalability analysis, however data might be loaded as a parametric study. For example, in the following example, the data has been loaded with three problem sizes, MIN, HALF and FULL.

0 0 PerfExplorer Client File Analysis Views Charts Visualization Help Analysis Management Q Cluster Results Correlation Results ► D SHAMRC ► 3 SMG2000 Field Value SPhot Name HPM016 Experiment ID 🕨 🧊 Uintah system_name ► 🔰 WRE system_machine_type ▶ jjj gyro.81-std system_arch ▼ 河 gyro.B1~std.HPM system_os ▼ D HPM018 system_memory_size system_processor_amt ▶ [] FULL 1.6 system_I1_cache_size ► III HALF16 system_l2_cache_size ► [] MIN16 system_userdata ▼ I HPM032 compiler_cpp_name ► 📁 FULL32 compiler_cpp_version compiler_cc_name ► | | HALF32 compiler cc version ▶ [MIN32 compiler_java_dirpath 📝 河 НРМО64 compiler_lava_version ▶ 3 FULL 64 compiler_userdata ► III HALF84 configure_prefix ► | MIN64 configure_arch configure_cpp configure_cc ► III FULL98 configure_jdk ► 3 HALF96 configure_profile ► [] MIN96 configure_userdata HPM128 userdata ▶ | FULL128 ► 3 HAUF128 ▶ | MIN128

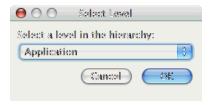
Figure 9.1. Potential scalability data organized as a parametric study

In order to examine this data in a scalability study, it is necessary to reorganize the data. However, it is not necessary to re-load the data. Using views in PerfExplorer, you can re-organize the data based on values in the database.

9.1. Creating Views

To create a view, select the "Create New View" item under the "Views" main menu item. The first step is to select the table which will form the basis of the view. The three possible values are Application, Experiment and Trial:

Figure 9.2. Selecting a table



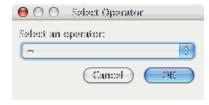
After selecting the table, you need to select the column on which to filter:

Figure 9.3. Selecting a column



After selecting the column, you need to select the operator for comparing to that column:

Figure 9.4. Selecting an operator



After selecting the operator, you need to select the value for comparing to the column:

Figure 9.5. Selecting a value



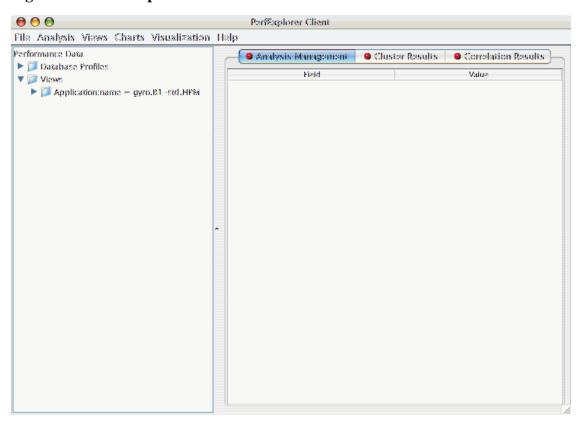
After selecting the value, you need to select a name for the view:

Figure 9.6. Entering a name for the view



After creating the view, you will need to exit PerfExplorer and re-start it to see the view. This is a known problem with the application, and will be fixed in a future release.

Figure 9.7. The completed view



9.2. Creating Subviews

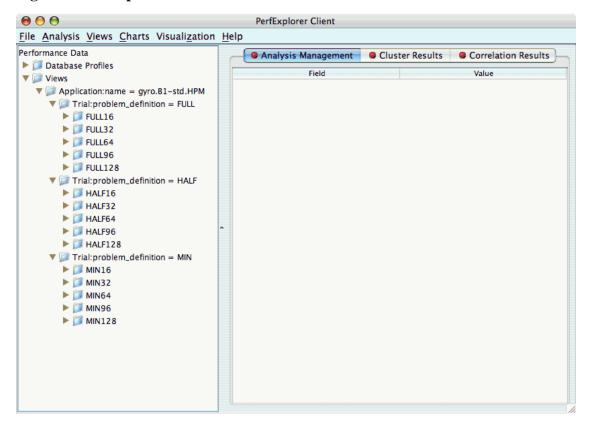
In order to create sub-views, you first need to select the "Create New Sub-View" item from the "Views" main menu item. The first dialog box will prompt you to select the view (or sub-view) to base the new sub-view on:

Figure 9.8. Selecting the base view



After selecting the base view or sub-view, the options for creating the new sub-view are the same as creating a new view. After creating the sub-view, you will need to exit PerfExplorer and re-start it to see the sub-view. This is a known problem with the application, and will be fixed in a future release.

Figure 9.9. Completed sub-views



Chapter 10. Running PerfExplorer Scripts

As of version 2.0, PerfExplorer has officially supported a scripting interface. The scripting interface is useful for adding automation to PerfExplorer. For example, a user can load a trial, perform data reduction, extract out key phases, derive metrics, and plot the result.

10.1. Analysis Components

There are many operations available, including:

- BasicStatisticsOperation
- CopyOperation
- CorrelateEventsWithMetadata
- CorrelationOperation
- DeriveMetricOperation
- DifferenceMetadataOperation
- DifferenceOperation
- DrawBoxChartGraph
- DrawGraph
- DrawMMMGraph
- ExtractCallpathEventOperation
- ExtractEventOperation
- ExtractMetricOperation
- ExtractNonCallpathEventOperation
- ExtractPhasesOperation
- ExtractRankOperation
- KMeansOperation
- LinearRegressionOperation
- LogarithmicOperation
- MergeTrialsOperation
- MetadataClusterOperation
- PCAOperation

- RatioOperation
- ScalabilityOperation
- TopXEvents
- TopXPercentEvents

10.2. Scripting Interface

The scripting interface is in Python, and scripts can be used to build analysis workflows. The Python scripts control the Java classes in the application through the Jython interpreter (http://www.jython.org/). There are two types of components which are useful in building analysis scripts. The first type is the PerformanceResult interface, and the second is the PerformanceAnalysisComponent interface. For documentation on how to use the Java classes, see the javadoc in the perfexplorer source distribution, and the example scripts below. To build the perfexplorer javadoc, type

```
%>./make javadoc
```

in the perfexplorer source directory.

10.3. Example Script

```
from glue import PerformanceResult
from glue import PerformanceAnalysisOperation
from glue import ExtractEventOperation
from glue import Utilities
from glue import BasicStatisticsOperation
from glue import DeriveMetricOperation
from glue import MergeTrialsOperation
from glue import TrialResult
from glue import AbstractResult
from glue import DrawMMMGraph
from edu.uoregon.tau.perfdmf import Trial
from java.util import HashSet
from java.util import ArrayList
True = 1
False = 0
def glue():
        print "doing phase test for gtc on jaquar"
        # load the trial
        Utilities.setSession("perfdmf.demo")
        trial1 = Utilities.getTrial("gtc_bench", "Jaguar Compiler Options", "fasts
        result1 = TrialResult(trial1)
        print "got the data"
        # get the iteration inclusive totals
        events = ArrayList()
        for event in result1.getEvents():
                #if event.find("Iteration") >= 0 and result1.getEventGroupName(eve
                if event.find("Iteration") >= 0 and event.find("=>") < 0:</pre>
                        events.add(event)
```

extractor = ExtractEventOperation(result1, events)

```
extracted = extractor.processData().get(0)
print "extracted phases"
# derive metrics
derivor = DeriveMetricOperation(extracted, "PAPI L1 TCA", "PAPI L1 TCM", D
derived = derivor.processData().get(0)
merger = MergeTrialsOperation(extracted)
merger.addInput(derived)
extracted = merger.processData().get(0)
derivor = DeriveMetricOperation(extracted, "PAPI_L1_TCA-PAPI_L1_TCM", "PAP
derived = derivor.processData().get(0)
merger = MergeTrialsOperation(extracted)
merger.addInput(derived)
extracted = merger.processData().get(0)
derivor = DeriveMetricOperation(extracted, "PAPI_L1_TCM", "PAPI_L2_TCM", D
derived = derivor.processData().get(0)
merger = MergeTrialsOperation(extracted)
merger.addInput(derived)
extracted = merger.processData().get(0)
derivor = DeriveMetricOperation(extracted, "PAPI L1 TCM-PAPI L2 TCM", "PAP
derived = derivor.processData().get(0)
merger = MergeTrialsOperation(extracted)
merger.addInput(derived)
extracted = merger.processData().get(0)
derivor = DeriveMetricOperation(extracted, "PAPI FP INS", "P WALL CLOCK TI
derived = derivor.processData().get(0)
merger = MergeTrialsOperation(extracted)
merger.addInput(derived)
extracted = merger.processData().get(0)
derivor = DeriveMetricOperation(extracted, "PAPI_FP_INS", "PAPI_TOT_INS",
derived = derivor.processData().get(0)
merger = MergeTrialsOperation(extracted)
merger.addInput(derived)
extracted = merger.processData().get(0)
print "derived metrics..."
# get the Statistics
dostats = BasicStatisticsOperation(extracted, False)
stats = dostats.processData()
print "got stats..."
return
for metric in stats.get(0).getMetrics():
        grapher = DrawMMMGraph(stats)
        metrics = HashSet()
        metrics.add(metric)
        grapher.set_metrics(metrics)
        grapher.setTitle("GTC Phase Breakdown: " + metric)
        grapher.setSeriesType(DrawMMMGraph.TRIALNAME);
        grapher.setCategoryType(DrawMMMGraph.EVENTNAME)
        grapher.setValueType(AbstractResult.INCLUSIVE)
        grapher.setXAxisLabel("Iteration")
grapher.setYAxisLabel("Inclusive " + metric);
        # grapher.setLogYAxis(True)
        grapher.processData()
# graph the significant events in the iteration
subsetevents = ArrayList()
```

```
subsetevents.add("CHARGEI")
        subsetevents.add("PUSHI")
        subsetevents.add("SHIFTI")
        print "got data..."
        for subsetevent in subsetevents:
               events = ArrayList()
               for event in result1.getEvents():
                       if event.find("Iteration") >= 0 and event.rfind(subseteven
                               events.add(event)
               extractor = ExtractEventOperation(result1, events)
               extracted = extractor.processData().get(0)
               print "extracted phases..."
               # get the Statistics
               dostats = BasicStatisticsOperation(extracted, False)
               stats = dostats.processData()
               print "got stats..."
                for metric in stats.get(0).getMetrics():
                       grapher = DrawMMMGraph(stats)
                       metrics = HashSet()
                       metrics.add(metric)
                       grapher.set_metrics(metrics)
                       grapher.setTitle(subsetevent + ", " + metric)
                       grapher.setSeriesType(DrawMMMGraph.TRIALNAME);
                       grapher.setCategoryType(DrawMMMGraph.EVENTNAME)
                       grapher.setValueType(AbstractResult.INCLUSIVE)
                       # grapher.setLogYAxis(True)
                       grapher.processData()
        return
print "-----" JPython test script start -----"
glue()
print "----- JPython test script end -----"
```

Chapter 11. Derived Metrics

Sometimes metrics in a profile need to be combined to create a derived metric. PerfExplorer allows the user to create these using the derived metric expression tab.

11.1. Creating Expressions

The text box at the top of the tab allows the user to enter an expression. Double clicking on a metric in the "Performance Data" tree will copy that metrics name into the box. If a metric contains any operands, the whole metric must be surrounded by quotes. If the you would like of the metric to be renamed, then you should start the expression with the new name and and equals sign.

If this is the only metric you wish to derive, then select the trial, expression or application where the metric should be derived and then click apply. If you wish to derive many metrics, then click Add to List and create more expressions.

11.2. Selecting Expressions

If you have added multiple expressions, you can select one or many of them to apply. They will be derived from top to bottom. After you have select some, you can select the trial, experiment or application to apply the expression to and then click apply.

11.3. Expression Files

You can also derive metrics using an expression file. An expression file has a single expression per line. To parse the file, select the trial, experiment or application to apply the expressions to; then select File > Parse Expression File and chose the file.